

862750-2750760

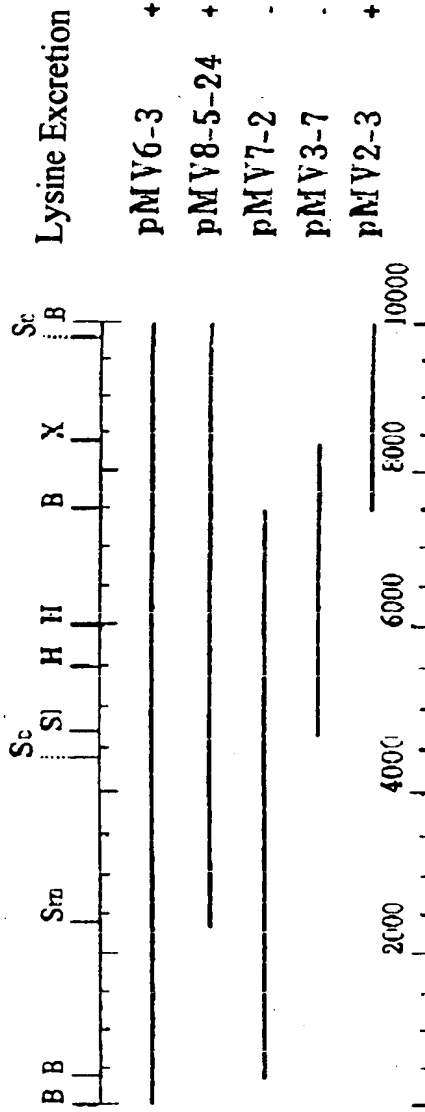


Figure 1

CgLyse 1 MVIMEIFITGLLLGASLLLSIGPONVLVIKOGIKREGLIAVLLVCI.TSDV 50  
 EcYgga 1 .....MILPLGPQNAFVMNOGIRROYHIMIALLCALSDL 34  
  
 CgLyse 51 ELFIAGTLGVDLLSNAAPIVLDIMRWGGIAYLLWFAVMAAKDAMTNKVEA 100  
 EcYgga 35 VLICAGIFGGSALLMQSPWLLALVTWGGVAFLLWYGFAGFTAMSSNIE 83  
  
 CgLyse 101 PQIIETEPTVPDDTPLGGSAVATDTRNRVRVEVSVDKQORVWVKPMLMAI 150  
 EcYgga 84 .....LASAEVMKQGRWK.....IIATMLAV 104  
  
 CgLyse 151 VTWLNPNAYLDAFVFIGGVGAQYGDTRWIFACAFASLIWFPLVGEG 200  
 EcYgga 105 ..TWLNPHVYLDTEVVLGSLGGQLDVEPKRWFAFGTISASFLWFFGLALL 152  
  
 CgLyse 201 AAALSRLSSPKVWRWINVVAVVMTALAIKMLMG..... 236  
 EcYgga 153 AAWLAPRLRTAKAQRIINLVVGCVMWFIALQLARDGIAHAQALFS 197

Figure 2

# Complementation of Export defects

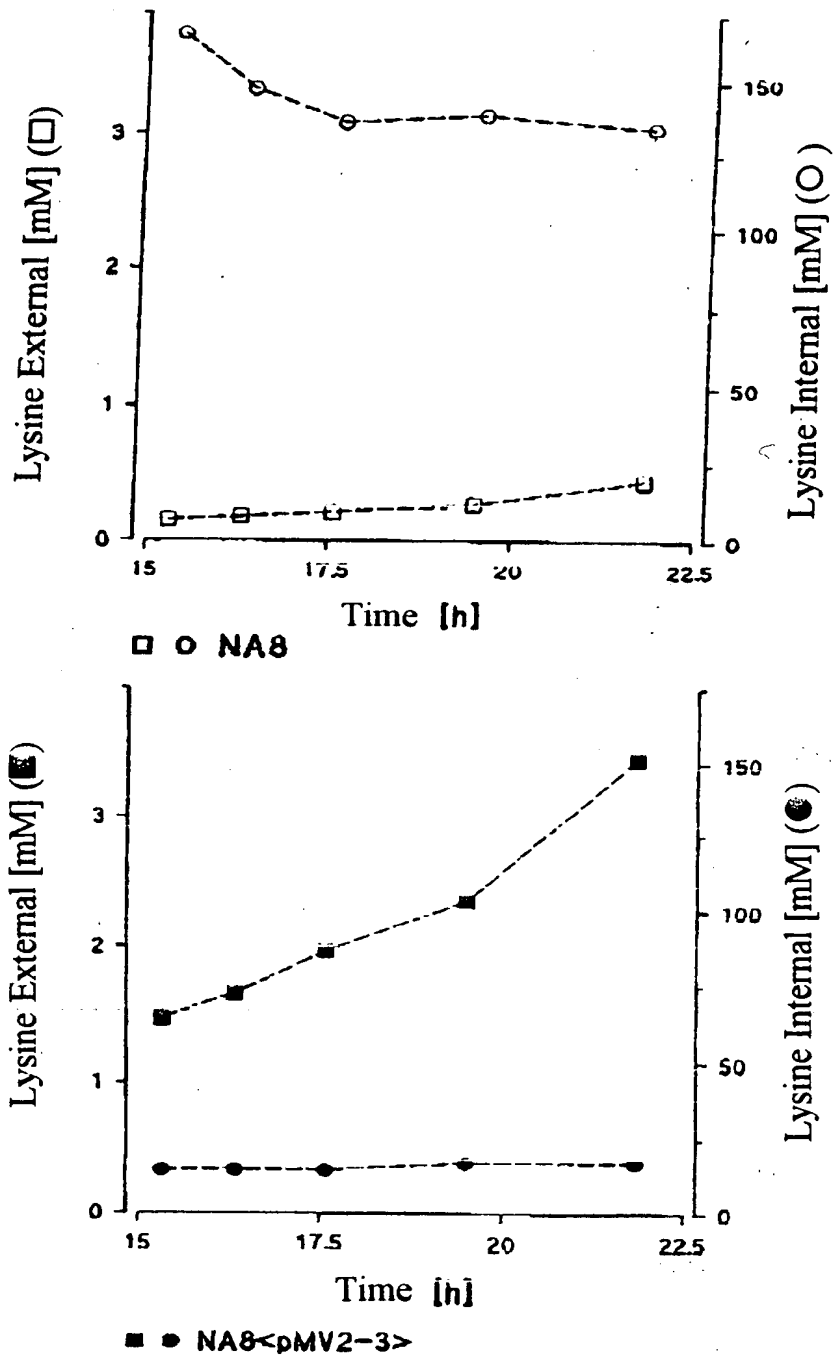


Figure 3

202790-2750160

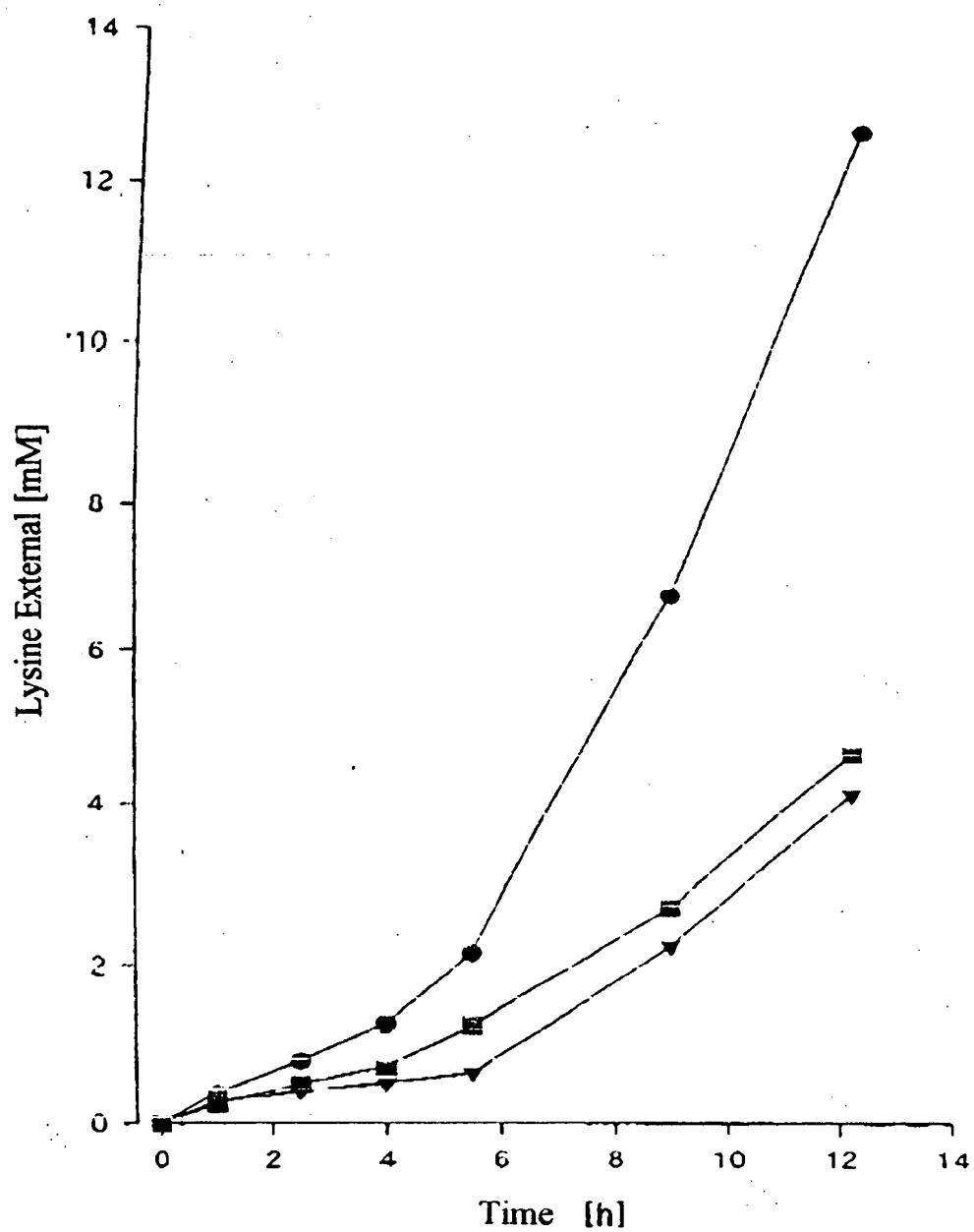


Figure 4